60

-10297.204.ST25.txt SEQUENCE LISTING Novozymes A/S <110> Jørgensen, Steen Troels Rasmussen, Michael Dolberg Andersen, Jens Tønne Olesen, Peter Bjarke Clausen, Ib Groth Improved Bacillus Host Cell <120> 10297.204 <130> <160> 22 PatentIn version 3.2 <170> <210> <211> 1602 <212> DNA Bacillus licheniformis <213> <220> <221> CDS (501)..(1421) <400> aacagccaga aaggagacga agaacgtctc tatttaacag agcttgatat gttctgcgtc - 60 catttcgggt caaggatata ttcactatta tgaaaccaat ttttccgatt tccgtataaa 120 caaagcgtgg aacctgttaa aggacaagaa aaatcgacag ggatataaaa atcggcaagc 180 tttccaaagg gaaccggggc aggctgaagc tagcgctcac attggcgagc gatgcagcgg 240 300 tcgtgctgtt tgatgaacca tccggtctcc gatggtgagg gagtcgacag tcaaaagcct tcttacatat ttggagcttg acaagcagac aatcgccatc gccacctatg aaattgaaca 360 420 aatattggat gtcgtcttaa atggccgaat tgaagcaaaa aaggatgtcg aacagctgcg cgaggaatcc ggaatgtcgc ttttggaatg gctgaaacgg tttattaaag aaaaaaattg 480 aaaaagcggg aggagaaaca ttg gaa aca ctt ttg gaa tta aaa aat gta tca 533 Leu Glu Thr Leu Leu Glu Leu Lys Asn Val Ser aaa acg atc agg ggg aaa aag atc atc gag ggc ttg agt ttt gac gtg Lys Thr Ile Arg Gly Lys Lys Ile Ile Glu Gly Leu Ser Phe Asp Val 15 20 25 581 cgg gca ggc gag ata ttc ggc ttc ctg ggg ccg aac ggc gcc gga aaa Arg Ala Gly Glu Ile Phe Gly Phe Leu Gly Pro Asn Gly Ala Gly Lys 30 35 40 629 acg acg acg atc cgg atg att gtc gga cat atg agc att acc gcc ggt Thr Thr Ile Arg Met Ile Val Gly His Met ser Ile Thr Ala Gly 677 gaa atc gcc gtg tgc ggc gtc agc gta aaa gaa aac ttt gaa aag gct Glu Ile Ala Val Cys Gly Val Ser Val Lys Glu Asn Phe Glu Lys Ala 725

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Leu Ser Thr Leu Met Phe Glu Gly Asp Ala Lys Glu Leu Thr Leu Thr
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Page 4

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Ser Leu Leu Thr Ala Ser Ile Ala Ala Leu Glu Lys Leu Lys Glu Tyr 65 70 75 80

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Bacillus licheniformis

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Leu		Arg 110	Pro	۷al	Asp	Leu		L0297 Arg					Thr	Glu	Αla	
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Leu Asp Leu Ile Leu Ala Glu Glu Thr Ala Tyr Arg Ala Ser Glu Gln 180 185 190

Tyr Gln Glu Asp Arg Arg Phe Trp Leu Gly Arg Phe Ser Asp Glu Pro 195 200 205

Glu Val Val Ser Leu Ala Glu Arg Ala Pro Arg Thr Ser Ser Phe 210 215 220

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10297.204.ST25.txt 250

245

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10297.204.ST25.txt 520 525

515

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10297.204.sT25.txt 795

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10297.204.ST25.txt 1570 157

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10297.204.ST25.txt 1830

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10297.204.ST25.txt 2075 2080 2085

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PCT/DK03/00199

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Thr Asp Ala Ser Pro Ile Ile Val Tyr Gly His Met Lys Pro Val Met 50 60

Ala Val Ser Phe Leu Ala Cys Val Lys Ala Gly His Pro Tyr Ile Pro 65 70 75 80

Val Asp Val Ser Ile Pro Ala Asp Arg Ile Leu L<u>y</u>s Ile Ile Asn Ser 85 90 95

Ser Lys Ala Glu Leu Leu Leu Asn Asn Ser Gly Thr Ser Val Asp Thr 100 105 110

Gly Asp Ala Leu Ile Ser Val Val Glu Pro Gly Val Leu Glu Gly Asp 115 120 125

Gly Val Pro Glu Thr Asp Pro Gly Arg Trp Val His Gly Glu Asp Thr 130 135 140

Phe Tyr Ile Ile Tyr Thr Ser Gly Ser Thr Gly Asn Pro Lys Gly Val 145 150 155 160

Gln Ile Ser Ala Asp Asn Leu Gln Ser Phe Thr Asp Trp Ile Thr Asn 165 170 175

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Ser Phe Asp Leu Ser Val Met Asp Leu Tyr Pro Cys Leu Gln Ser Gly
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Gly Thr Leu Trp Thr Val Thr Lys Asp Met Ile Asn Arg Pro Lys Leu 210 215 220

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Leu Leu Pro Glu Leu Ser Leu Phe Met Phe Cys Gly Glu Thr Leu Pro Page 35 10297.204.ST25.txt 265

260

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Glu Val Thr Asp Asp Val Leu Asn Lys Tyr Ser Ser Leu Pro Val Gly 305 310 315

Ser Glu Lys Pro Glu Thr Glu Ile Val Ile Ile Asn Glu Asp Gly Lys 325 330 335

Ala Val Gln Asp Gly Glu Lys Gly Glu Ile Ile Ile Thr Gly Ala Ser

val Ser Lys Gly Tyr Leu Gly Glu Lys Ala Leu Thr Glu Lys Ala Phe 355 360 365

Phe Ser Tyr Asn Gly Ser Pro Ala Tyr Arg Thr Gly Asp Ala Gly Tyr 370 375 380

Lys Glu Asn Gly Gln Leu Phe Phe Leu Gly Arg Leu Asp Phe Gln Ile 385 390 395 400

Lys Leu His Gly Tyr Arg Ile Glu Leu Glu Glu Ile Glu Tyr Gln Ile 405 410 415

Asn Gln Ser Arg Tyr Val Gln Ser Ala Val Val Ile Pro Phe Tyr Arg 420 425 430

Glu Glu Lys Ile Glu Tyr Leu Ile Ala Met Ile Val Pro Ala Glu His 435 440 445

Asp Phe Glu Lys Glu Tyr Gln Leu Thr Ser Ala Ile Lys Lys Asp Leu 450 455 460

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10297.204.ST25.txt 445 450 455

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ggg Gly	gat Asp 845	act Thr	gca Ala	cgt Arg	tgg Trp	ctt Leu 850	cct Pro	gac Asp	ggt Gly	aca Thr	gtt Val 855	gat Asp	ttt Phe	atc Ile	ggc Gly	3077	
cga Arg 860	aga Arg	gac Asp	gat Asp	caa Gln	gtg Val 865	aaa Lys	atc Ile	cgc Arg	ggc Gly	ttc Phe 870	cgg Arg	atc Ile	gag Glu	ctt Leu	ggc Gly 875	3125	·.
gag Glu	ata Ile	gag Glu	tcg Ser	gtg Val 880	ctg Leu	caa Gln	gga Gly	gcc Ala	cct Pro 885	gct Ala	gtc Val	gaa Glu	aaa Lys	gca Ala 890	vaı	3173	
gtc Val	ctc Leu	gcg Ala	cgg Arg 895	cat His	gaa Glu	acc Thr	ggc Gly	ggc Gly 900	agc Ser	ctt Leu	gag Glu	gtt Val	tgc Cys 905	gct Ala	tat Tyr	3221	
gtc Val	gtt Val	ccg Pro 910	aaa Lys	caa Gln	ggc Gly	gga Gly	aag Lys 915	att Ile	cac His	att Ile	cag Gln	gga Gly 920	ctg Leu	agg Arg	gag Glu	3269	, je salikari
cac His	tta Leu 925	tcc Ser	aaa Lys	cac His	ctg Leu	cct Pro 930	Asp	tat Tyr	atg Met	atc Ile	cct Pro 935	Ser	tgc Cys	ttt Phe	gtc Val	3317	
gag Glu 940	ctt Leu	aat Asn	gaa Glu	ata Ile	ccg Pro 945	Leu	acg Thr	gca Ala	agc Ser	gga Gly 950	Lys	gtc Val	gac Asp	cgg Arg	aaa Lys 955	3365	
gcg Ala	ctg Leu	ctt Leu	cgc Arg	cat His 960	Glu	gtg Val	tct Ser	gtc Val	agc Ser 965	Gly	act Thr	gca Ala	gaa Glu	tat Tyr 970	gcg Ala	3413	
Ālā	Pro	Arg	Asn 975	Glu	Cys	Glu	Glu	980	Met	\ Val	Gly	lle	985	GIR	gaa Glu	3461	
gtg Val	ctc Leu	gga Gly	gct Ala	gaa Glu	caa Gln	gtc Val	gga Gly	atc Ile	H15	gac Asp ge 4	GIN	ttt Phe	tt Ph	t ga ie As	t ctg p Leu	3509	

10297.204.ST25.txt 990 995 1000

	9	190				95	כי				TOC				
ggc Gly	ggc Gly 1005	cac His	tct Ser	ctc Leu	aaa Lys	gcg Ala 1010	atg Met	acc Thr	atg Met	ctg Leu	gca Ala 1015	aaa Lys	att Ile	cat His	3554
aaa Lys	gct Ala 1020	ttc Phe	gga Gly	gtc Val	gaa Glu	gtt Val 1025	ccg Pro	ttg Leu	caa Gln	gtc Val	cta Leu 1030	ttt Phe	gaa Glu	aag Lys	3599
	aca Thr 1035	gta Val	gcg Ala	gcg Ala	ctg Leu	agc Ser 1040	ggc Gly	ttt Phe	gtt Val	tct Ser	gaa Glu 1045	gcc Ala	gag Glu	aaa Lys	3644
_	gga Gly 1050	ttc Phe	gcg Ala	gta Val	att Ile	gaa Glu 1055	ccg Pro	gcg Ala	ccg Pro	gaa Glu	agt Ser 1060	gac Asp	gat Asp	tat Tyr	3689
	ctt Leu 1065	tca Ser	ttg Leu	gcg Ala	cag Gln	cag Gln 1070	cga Arg	atc Ile	tat Tyr	atc Ile	gtc Val 1075	agc Ser	cag Gln	ctg Leu	3734
	ggt Gly 1080	gtc Val	ggc Gly	tac Tyr	aat Asn	atg Met 1085	ccg Pro	gcc Ala	gcg Ala	gcg Ala	atg Met 1090	ctt Leu	gaa Glu	ggc Gly	3779
	ctt Leu 1095	gac Asp	tct Ser	ggg Gly	cgc Arg	ctt Leu 1100	gaa Glu	gct Ala	gct Ala	ttt Phe	caa Gln 1105	aaa Lys	ctg Leu	att Ile	3824
	agg Arg 1110	cat His	gag Glu	gcg Ala	ctg Leu	cgt Arg 1115	act Thr	tcg Ser	ttt Phe	aca Thr	gtt Val 1120	gtg Val	gat Asp	ggc Gly	3869
gag Glu	ccg Pro 1125	agg Arg	cag Gln	acc Thr	gtt val	cat His 1130	caa Gln	agg Arg	gtt Val	cag Gln	ttc Phe 1135	aag Lys	att Ile	gag Glu	39 <u>1</u> 4
_	gtt Val 1140	Lys	gcg Ala	gaa Glu	ggc Gly	aag Lys 1145	ccg Pro	atc Ile	gaa Glu	caa Gln	atc Ile 1150	gcg Ala	aaa Lys	tca Ser	3959
ttt Phe	gtc Val 1155	Arg	cgc Arg	ttt Phe	gat Asp	ctc Leu 1160	gcc Ala	aaa Lys	gct Ala	ccg Pro	ctg Leu 1165	atg Met		gcg Ala	4004
ggt Gly	ctc Leu 1170	Val	agc Ser	ctt Leu	Ala	gac Asp 1175	Gly	aga Arg	cat His	ctg Leu	ctg Leu 1180	Leu	ttc Phe	gat Asp	4049
atg Met	cat His 1185	cat His	ctc Leu	gta Val	tct Ser	gac Asp 1190	ggc Gly	gtg Val	tcg Ser	att Ile	tcg Ser 1195	atc Ile	ata Ile	ttg Leu	4094
aat Asn	gaa Glu 1200	Leu	gcc Ala	gcg Ala	ctg Leu	tat Tyr 1205	Asn	ggg Gly	gag Glu -	gaa Glu	ttg Leu 1210	Pro	gag Glu	ctg Leu	4139
cgt Arg	ttg Leu 1215	His	tac Tyr	aaa Lys	gat Asp	tat Tyr 1220	ÃΊa	ttc Phe	tgg Trp	caa Gln	agg Arg 1225	Ālā	cag Gln	gca Ala	4184
caa Gln	gaa Glu 1230	Gly	ttt Phe	caa Gln	aaa Lys	gaa Glu 1235	Glu	gcc Ala	tat Tyr	tgg Trp	gaa Glu 1240	ser		ttt Phe	4229
	ggc Gly					ctg Leu	cag Gln	Leu	atg Met Page	Thr	gac Asp	gag Glu	ccg Pro	agg Arg	4274
	_														

	1245		,			1250	102	297.2	204.5	T25	txt 1255					
ccg Pro		gtt Val	caa Gln	agc Ser	ttt Phe	gaa Glu 1265	gga Gly	gac Asp	agg Arg	gta Val	tcc ser 1270	gca Ala	gtc Val	ctg Leu	•	4319
	aaa Lys 1275	gac Asp	ctg Leu	aaa Lys	gaa Glu	aaa Lys 1280	tta Leu	gcc Ala	gtg Val	ctg Leu	gca Ala 1285	gag Glu	caa Gln	aac Asn		4364
ggc	gcg Ala 1290	aca Thr	ttg Leu	tac Tyr	atg Met	gtc Val 1295	atg Met	ctg Leu	tct Ser	gcg Ala	tac Tyr 1300	aac Asn	atg Met	ctt Leu		4409
ctt Leu	gca Ala 1305	aag Lys	tac Tyr	tcc Ser	ggg Gly	cag Gln 1310	gaa Glu	gat Asp	gtg Val	atc Ile	gtc Val 1315	ggg Gly	acg Thr	ccg Pro		4454
gcg Ala	gcc Ala 1320	ggc Gly	cgc Arg	aga Arg	cat His	tcc Ser 1325	gat Asp	ctg Leu	gaa Glu	gga Gly	atc Ile 1330	atc Ile	ggc Gly	atg Met		4499
ttt Phe	gtc Val 1335	aat Asn	acg Thr	ctg Leu	gcg Ala	atc Ile 1340	cgt Arg	tca Ser	aaa Lys	gtg Val	gac Asp 1345	ccg Pro	ggc Gly		•	4544
acg Thr	ttc Phe 1350	Ā٦a	gac Asp	ttc Phe	ctg Leu	aat Asn 1355	gat Asp	gta Val	aag Lys	aaa Lys	acg Thr 1360	gta Val	atc Ile	gat Asp		4589
gct Ala	ttt Phe 1365	gag Glu	cat His	cag Gln	gat Asp	tat Tyr 1370	ccg Pro	ttt Phe	gag Glu	cgg Arg	ctt Leu 1375	gcg Ala	gag Glu	aag Lys		4634
ttt Phe	ggc Gly 1380	Ala	tcg Ser	cgc Arg	gat Asp	tta Leu 1385	agc Ser	cgt Arg	cat His	CCC Pro	atc Ile 1390	rne	gac Asp	acg Thr		4679
	ttt Phe 1395	Ile	ttg Leu	cag Gln	aat Asn	gca Ala 1400	tgg Trp	gaa Glu	gac Asp	att Ile	ccg Pro 1405	Leu	ctc Leu	ggc Gly		4724
Asp	ttg Leu 1410	His	ctc Leu	tcg Ser	att Ile	tat Tyr 1415	gaa Glu	acg Thr	aac Asn	ttc Phe	aat Asn 1420	Tie	gca Ala	aaa Lys		4769
ttc Phe	gac Asp 1425		acg Thr	ctg Leu	cag Gln	gca Ala 1430	aaa Lys	gaa Glu	gag Glu	cag Gln	gga Gly 1435	gaa Glu	ttg Leu	ata Ile		4814
ctc Leu	gat Asp 1440	ttg Leu	gat Asp	tac Tyr	agc Ser	acg Thr 1445	aag Lys	ctg Leu	ttt Phe	aaa Lys	aaa Lys 1450	gat Asp	acg Thr	gct Ala		4859
	agg Arg 1455	Met	ctg Leu	aaa Lys	gca Ala	tac Tyr 1460	Leu	aac Asn	ctt Leu	ctg Leu	gag Glu 1465	Asp	atg Met	gcg Ala		4904
	gat Asp 1470	Pro	atg Met	ctc Leu	agg Arg	atc Ile 1475	Gly	gag	tac Tyr	ago Ser	ctt Leu 1480	Leu	aca Thr	gaa Glu		4949
	gaa Glu 1485	Thr	aat Asn	cgc Arg	cag Gln	ctc Leu 1490	٧a٦	gcg Ala	tto Phe	aac Asr	ccc Pro 1495	Ala	aca Thr	agc Ser		4994
gat Asp	tat Tyr	ccg Pro	g cgg Arg	gag Glu	aaa Lys	acg Thr	atc Ile	gtt Val	cag Glr Page	cta Leu 42	ttc Phe	gag	gaa Glu	cag Gln		5039

	1500					1505	102	97.2	204.5	т25.	txt 1510				
gct Ala	gca Ala 1515	gaa Glu	aga Arg	ggc Gly	ggt Gly	tat Tyr 1520	ccc Pro	gcc Ala	ctg Leu	caa Gln	ttt Phe 1525	gaa Glu	gac Asp	aaa Lys	5084
gta Val	tgg Trp 1530	tca Ser	tat Tyr	gat Asp	gag Glu	ctc Leu 1535	aat Asn	cgg Arg	aaa Lys	gcg Ala	aac Asn 1540	caa Gln	ttg Leu	gca Ala	5129
	cgc Arg 1545	ctc Leu	cgc Arg	gaa Glu	agt Ser	ggc Gly 1550	gtg Val	caa Gln	gcc Ala	ggc Gly	acg Thr 1555	acg Thr	gcg Ala	gcc Ala	5174
att Ile	ctt Leu 1560	acg Thr	gcc Ala	agg Arg	tcg Ser	gct Ala 1565	gaa Glu	atg Met	gtc Val	atc Ile	ggc Gly 1570	att Ile	ctc Leu	gct Ala	52 <u>1</u> 9
	ctg Leu 1575	aaa Lys	gcc Ala	ggc Gly	ggg Gly	gct Ala 1580	tat Tyr	gtg Val	ccg Pro	atc Ile	gat Asp 1585	ccc Pro	gat Asp	cat His	5264
ccg Pro	gaa Glu 1590	aag Lys	cgt Arg	gtg Val	caa Gln	cat His 1595	ttc Phe	ttc Phe	aag Lys	gac Asp	agc Ser 1600	gga Gly	gcg Ala	gcg Ala ,	5 30 9
gtt Val	ctt Leu 1605	tta Leu	acg Thr	caa Gln	aag Lys	gcg Ala 1610	atg Met	aag Lys	ccg Pro	ctc Leu	gct Ala 1615	gag Glu	gcg Ala	gcg Ala	5354
gaa Glu	ttt Phe 1620	Gly	ggc Gly	gac Asp	atc Ile	cta Leu 1625	ttt Phe	gtt Val	gag Glu	gat Asp	gaa Glu 1630	aac Asn	ctt Leu	tat Tyr	5399
atg Met	gga Gly 1635	Asp	gcg Ala	tct Ser	gat Asp	ttg Leu 1640	cgc Arg	ctt Leu	ccg Pro	att Ile	tct Ser 1645	ccg Pro	gaa Glu	gca Ala	5444
atg Met	gca Ala 1650	Asn	cta Leu	acg Thr	tac Tyr	aca Thr 1655	tct Ser	gga Gly	acg Thr	aca Thr	ggg Gly 1660	Thr		aag Lys	5489
	aat Asn 1665	Met	gtc Val	tct Ser	cac His	cgg Arg 1670	Asn	att Ile	ttg Leu	cgt Arg	act Thr 1675	Val	aaa Lys	aac Asn	5534
gcg Ala	aat Asn 1680	tac Tyr	ttg Leu	gaa Glu	gtg Val	atg Met 1685	gaa Glu	tcc Ser	gat Asp	atc Ile	gtt Val 1690	ctt Leu	tcg Ser	att Ile	5579
	aat Asn 1695	Tyr	gtg Val	ttc Phe	gac Asp	gcc Ala 1700	Phe	atg Met	ttt Phe	gat Asp	gtg Val 1705	Phe		tca Ser	5624
ttg Leu	ctg Leu 1710		ggc	gcc Ala	aag Lys	ctg Leu 1715	gtg Val	ata Ile	gct Ala	cca Pro	aaa Lys 1720	gac Asp	acg Thr	att Ile	5669
	gat Asp 1725	Met	agc Ser	cgg Arg	ctc Leu	gca Ala 1730	His	gta Val	.Leu	gaa Glu	aaa Lys. 1735	GIU	aag Lys	att Ile	571
	att Tle 1740	Leu	atg Met	ato :Ile	acg Thr	acc Thr 1745	Ala	tta Leu	ttc Phe	aat Asr	ctg Leu 1750	Let	aca Thr	gac Asp	575
ato Met	g cgg Arg	ccg Pro	gac Asp	agt Ser	ttg Leu	aag Lys	gga Gly	/ Lei	cgc Arg Page	ı Arg	gtt Val	ct <u>c</u> Lei	tto Phe	ggc Gly	580

	1755					1760	102	97.2	204.5	T25	txt 1765					
ggc Gly	gaa Glu 1770	cgt Arg	gca Ala	tct Ser	gtc Val	gat Asp 1775	cat His	gtc Val	aga Arg	aga Arg	gcc Ala 1780	ctg Leu	aag Lys	aca Thr	58	49
gtt Val	ggc Gly 1785	aga Arg	ggg Gly	cgg Arg	ctg Leu	ctt Leu 1790	cat His	atg Met	tac Tyr	ggt Gly	ccg Pro 1795	tct Ser	gag Glu	agc Ser	58 ·	94
	gtc Val 1800	ttt Phe	aca Thr	acc Thr	tac Tyr	cat His 1805	cca Pro	gtc Val	aat Asn	gaa Glu			gac Asp		59	39
	cag Gln 1815	gct Ala	ata Ile	ccg Pro	atc Ile	gga Gly 1820	aag Lys	cct Pro	gtc Val	agc Ser	aat Asn 1825	acg Thr	gag Glu	gtt Val	59	84
ctg Leu	att Ile 1830	ctt Leu	gat Asp	tca Ser	ttc Phe	ggc Gly 1835	aat Asn	gta val	cag Gln	ccg Pro	gca Ala 1840	ggg Gly	gtt Val	gcg Ala	60	29
gga Gly	gaa Glu 1845	tta Leu	tgc Cys	gtc Val	ggc Gly	gga Gly 1850	gac Asp	ggc Gly	ctt Leu	gtc Val	cgc Arg 1855	ggc Gly	tat Tyr	ttt Phe	. 60	74
	cgc Arg 1860	cct Pro	gaa Glu	ctg Leu	acg Thr	gct Ala 1865	gaa Glu	aag Lys	ttt Phe	acg Thr	gcg Ala 1870	cat His	ccg Pro			L19
aaa Lys	acc Thr 1875	Gly	gaa Glu	aag Lys	att Ile	tac Tyr 1880	cgg Arg	acg Thr	ggt Gly	gat Asp	atg Met 1885	gcg Ala	aga Arg			L64
ctt Leu	gcc Ala 1890	Asp	gga Gly	tgc Cys	ctc Leu	gaa Glu 1895	ttt Phe	atc Ile	ggt Gly	cgg Arg	att Ile 1900	gat Asp	cat His	cag Gln		209
	aaa Lys 1905	Ile	cgc Arg	gga Gly	cag Gln	aga Arg 1910	atc Ile	gag Glu	ctg Leu	ggc Gly	gaa Glu 1915	atc Ile	gag Glu	cat His	62	254
His	Leu 1920	Leu	Thr	His	Asp	мет 1925	vaı	Gin	Giu	АІА	gct Ala 1930	vai	Leu	Ala	62	299
gtt Val	gac Asp 1935	acc Thr	ggg Gly	gcg Ala	gga Gly	gat Asp 1940	caa Gln	atg Met	atc Ile	tgt Cys	gca Ala 1945	tac Tyr	ttt Phe	acg Thr	63	344
	gat Asp 1950	G]n	gag Glu	ctt Leu	tcg Ser	agc Ser 1955	caa Gln	gag Glu	ctt Leu	cgc Arg	cga Arg 1960	HIS	gcg Ala	gca Ala	-	389
	ggt Gly 1965	Leu	cca Pro	ggc	tat Tyr	atg Met 1970	Ile	ccg Pro	tct Ser	gtg Val	Phe 1975	мет	cag Gln	ctt Leu	64	434
	gag Glu 1980	Leu				ggc Gly 1985	aat Asn	gga Gly	aag Lys	ato Ile	gac Asp 1990	Arg	agg Arg	gcg Ala	6-	479
	cct Pro 1995	Ğlı				gcg Ala 2000	Gln	gco Ala	gcg Ala	caa Glr	a aag n Lys 2005	Glu	tat Tyr	acg Thr	6	524
gcg Ala	cca Pro	agg Arg	ago Ser	ggc	aca Thr	gaa Glu	gcg Ala	. GIr	ctt Lei Page	Ala	gat Asp	ctt Leu	tgg Trp	cag Gln	. 6	569

	2010					2015	102	297.2	204.5	ST25.	txt 2020					
gag Glu	gtg Val 2025	ctc Leu	aat Asn	gta Val	ccg Pro	aaa Lys 2030	atc Ile	ggt Gly	gtg Val	cat His	gac Asp 2035	aat Asn	ttt Phe	ttt Phe		6614
	ctc Leu 2040	ggc Gly	gga Gly	cac His	tct Ser	tta Leu 2045	ctc Leu	gga Gly	atg Met	acg Thr	ctc Leu 2050	atc Ile	gcc Ala	cgc Arg		6659
	cag Gln 2055	caa Gln	gag Glu	atg Met	aat Asn	gtt Val 2060	gac Asp	ctg Leu	cag Gln	ctg Leu	aaa Lys 2065	gac Asp	ttg Leu	ttc Phe		6704
	gcg Ala 2070	ccg Pro	aca Thr	atc Ile	gaa Glu	agc Ser 2075	ctg Leu	gcc Ala	cag Gln	gcg Ala	gcg Ala 2080	gca Ala	aaa Lys	tct Ser		6749
	aaa Lys 2085	aaa Lys	tct Ser	gcc Ala	gtt Val	tat Tyr 2090	att Ile	gaa Glu	gcg Ala	gcg Ala	ccg Pro 2095	gac Asp	cgg Arg	gaa Glu		6794
acg Thr	tac Tyr 2100	ccc Pro	gtt Val	tcg ser	tcg Ser	gcg Ala 2105	caa Gln	aaa Lys	cgg Arg	ctg Leu	tac Tyr 2110	gtt Val	ctt Leu	cag Gln.		6839
cag Gln	att Ile 2115	gaa Glu	ggc Gly	gct Ala	gag Glu	aaa Lys 2120	agc Ser	tat Tyr	aac Asn	atg Met	ccg Pro 2125	gcc Ala	gtt Val	ctc Leu		6884
caa Gln	ctt Leu 2130	gaa Glu	gga Gly	aag Lys	ctt Leu	gac Asp 2135	ctg Leu	aag Lys	cgc Arg	ctg Leu	gag Glu 2140	tct Ser	gcc Ala	gcg Ala		6929
caa Gln	atg Met 2145	ctg Leu	atc Ile	aag Lys	cgc Arg	cat His 2150	gaa Glu	gcg Ala	ttc Phe	cgc Arg	acg Thr 2155	acg Thr	ttt Phe	gaa Glu		6974
	agg Arg 2160	Asn	ggc Gly	gaa Glu	cct Pro	gtt Val 2165	cag Gln	cgg Arg	att Ile	tgg Trp	gaa Glu 2170	Ala	gcg Ala	gaa Glu		7019
	acc Thr 2175	Ile	gat Asp	gtc Val	atc Ile	gat Asp 2180	Ala	gat Asp	gaa Glu	caa Gln	gaa Glu 2185	Ala	gaa Glu	aag Lys		7064
ctg Leu	att Ile 2190	Lys	gag Glu	ttc Phe	atc Ile	agg Arg 2195	ccg Pro	ttt Phe	gat Asp	tta Leu	aca Thr 2200	Lys	gct Ala	ccg Pro	•	7109
tta Leu	ttt Phe 2205	Arg	atg Met	agc Ser	atc Ile	ata Ile 2210	Arg	gtc Val	acg Thr	gaa Glu	gaa Glu 2215	Lys	cat His	ctt Leu		7154
	ctt Leu 2220	٧a٦	gat Asp	atg Met	cat His	cat His 2225	Ile	att	tca Ser	gac Asp	ggc Gly 2230	Ala	tcc Ser	gtc Val		7199
	gtg Val 2235	Leu	atc	gat Asp	gag Glu	atg Met 2240	Thr	agg Arg	ctt Leu	tat Tyr	gcg Ala 2245	Gly	gag Glu	gcg Ala		7244
	gaa Glu 2250	Pro	ctg Leu	agg Arg	att Ile	cag Gln 2255	Tyr	aaa Lys	gac Asp	tat Tyr	gct Ala 2260	٧a١	tgg Trp	cag Gln	•	7289
	cac His	ttg Leu	ctc Leu	act Thr	gaa Glu	cgg Arg	cac His	Lys	atg Met Page	Gin	gag Glu	gaa Glu	tac Tyr	tgg Trp		7334

	2265					2270	102	97.2	204.5	T25.	txt 2275					
	aag Lys 2280	gag Glu	ctt Leu	gac Asp	ggc Gly	gaa Glu 2285	ctg Leu	cct Pro	gtt Val	ttg Leu	acg Thr 2290	ctc Leu	ccg Pro	aca Thr	;	7379
	tat Tyr 2295	ccg Pro	cgt Arg	cct Pro	Ser	gtt Val 2300	caa Gln	acg Thr	ttt Phe	gaa Glu	gga G1y 2305	agc Ser	cgt Arg	atc Ile	;	7424
	ttt Phe 2310	tct Ser	ctg Leu	aaa Lys	ccg Pro	gag Glu 2315	ctt Leu	gtc Val	cag Gln	cag Gln	ctc Leu 2320	aga Arg	aga Arg	ctg Leu	•	7469
	aag Lys 2325	gaa Glu	acg Thr	gaa Glu	agc Ser	act Thr 2330	tta Leu	tac Tyr	atg Met	gtg Val	ctc Leu 2335	gct Ala	gcc Ala	tca Ser		7514
tac Tyr	ägt Ser 2340	acg Thr	ttt Phe	tta Leu	tcg Ser	aaa Lys 2345	ctg Leu	agt Ser	gga Gly	cag Gln	agc Ser 2350	gag Glu	gtc Val	att Ile		7559
ttc Phe	ggc Gly 2355	tct Ser	ccg Pro	gct Ala	gcc Ala	ggg Gly 2360	agg Arg	ccg Pro	cat His	gca Ala	gat Asp 2365	ttg Leu	agc Ser	cgt Arg		7604
atc Ile	atc Ile 2370	ggc Gly	atg Met	ttt Phe	gtc val	aat Asn 2375	acg Thr	ctt Leu	gcg Ala	ata Ile	aga Arg 2380	acg Thr	cgt Arg	cct Pro		7649
gaa Glu	gga Gly 2385	gac Asp	aag Lys	ccg Pro	ttt Phe	tct ser 2390	gca Ala	ttc Phe	ctt Leu	gag Glu	gaa Glu 2395	gta Val	aaa Lys	gaa Glu		7694
acg Thr	acg Thr 2400	Leu	ggc Gly	gca Ala	ttt Phe	gag Glu 2405	cat His	caa Gln	gat Asp	tac Tyr	ccg Pro 2410	Pne	gaa Glu	gaa Glu		7739
ctg Leu	atc Ile 2415	gag Glu	aag Lys	ctg Leu	aac Asn	atc Ile 2420	cag Gln	cgc Arg	gac Asp	atg Met	agc Ser 2425	Arg	aac Asn	ccg Pro		7784
tta Leu	ttt Phe 2430	Asp	gcc Ala	gtc Val	ttt Phe	tca ser 2435	atg Met	caa Gln	aat Asn	gcc Ala	gat Asp 2440	Leu	aaa Lys	gac Asp		7829
	tca Ser 2445	Met	gaa Glu	gga Gly	gtc Val	aca Thr 2450	ctt Leu	aaa Lys	ccg Pro	tat Tyr	gat Asp 2455	Pne	gcc Ala	cat His		7874
	act Thr 2460	Ala	aaa Lys	ttt Phe	gat Asp	ctg Leu 2465	Thr	ttg Leu	aca Thr	gcg Ala	gcg Ala 2470	Glu	gaa Glu	gac Asp		7919
	ctt Leu 2475	Leu	gtt Val	ttt Phe	gaa Glu	atg Met 2480	Glu	tac Tyr	aat Asn	aca Thr	gat Asp 2485	Leu	ttt Phe	aag Lys		7964
	gaa Glu 2490	ser				tgg Trp 2495	Ser	ggg Gly	tac Tyr	tgg Trp	gtc Val 2500	Asn	ctg Leu	ctc Leu		8009
gaa Glu	gcg Ala 2505	٧a٦	gcc Ala	gaa Glu	acc Thr	ccg Pro 2510	Asp	gca Ala	agt Ser	ctt Leu	tca Ser 2515	Asp	ctt Leu	tcg Ser		8054
ctt Leu	ctg Leu	gat Asp	gaa Glu	gcg Ala	gaa Glu	aaa Lys	cgg Arg	Arg	ato Ile Page	vai	caa Gln	aag Lys	tgg Trp	aat Asn		8099

	2520					2525	102	297.2	204.5	ST25	txt 2530					
	acg Thr 2535	aag Lys	ctc Leu	gct Ala	gtg Val	ccg Pro 2540	cag Gln	gac Asp	aaa Lys	acg Thr	gtc Val 2545	cat His	gag Glu	ctg Leu		8144
	gaa Glu 2550	gcg Ala	caa Gln	gtc Val	ctc Leu	cgc Arg 2555	act Thr	cct Pro	gac Asp	cgc Arg	ggg Gly 2560	gca Ala	gcg Ala	gtg Val		8189
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gca Ala	gag Glu 3330	gac Asp	aaa Lys	cgg Arg	atg Met	сас Ніѕ 3335	cat His	acg Thr	aaa Lys	acg Thr	gct Ala 3340	gag Glu	ttt Phe	tct Ser	10529
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11740

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 ~~~	~~~	<b>a</b> > >	cta	200	666	200	ast	c++	aas	ast

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Glu Ala Ile Arg Lys Phe Val Arg Thr Asn Asp Thr Met Arg Phe Arg 50 60

Leu Met Phe Glu Gly Glu Asp Glu Pro Lys Gln Tyr Ile Ala Glu Asp 65 70 75 80

Glu Pro Phe Gln Ile Glu Tyr Phe Asp Ala Ser Glu Ser Gly Gly Ala 85 90 95

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<213> Bacillus licheniformis

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Lys Lys Thr Val Ser Asp Gly Arg Ala Leu Ser Asp Tyr Tyr Arg Arg 675 680 685 His Gln Ile Asp Val Thr Asp Gly Thr Pro Ala His Leu Gln Leu Leu 690 700 Ala Ala Asp Asp Leu Ser Gly Val Lys Leu Arg His Met Leu Val 705 710 715 Gly Gly Glu Ala Leu Ser Arg Val Ala Thr Glu Arg Leu Leu Gln Leu 725 730 735 Phe Ala Glu Thr Ala Glu Ser Val Pro Ala Val Thr Asn Val Tyr Gly
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- Glu Val Ser Val Ser Gly Thr Ala Glu Tyr Ala Ala Pro Arg Asn Glu 965 970 975
- Cys Glu Glu Lys Met Val Gly Ile Trp Gln Glu Val Leu Gly Ala Glu 980 985 990
- Gln Val Gly Ile His Asp Gln Phe Phe Asp Leu Gly Gly His Ser Leu 995 1000 1005
- Lys Ala Met Thr Met Leu Ala Lys Ile His Lys Ala Phe Gly Val 1010 1020
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gga co Gly Pr	cg gac ro Asp 750	Arg	atc Ile	atc Ile	cat His	gta Val 755	tac Tyr	ggg Gly	ccg Pro	acg Thr	gaa Glu 760	acg Thr	acc Thr	gtc Val	2789
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whet Ille Glu Ser Asp 1005  cat tta ata gaa agc gat gac gac en loo yal leu loo yal leu loo yal loo yal loo yal leu loo yal loo yal leu loo yal	grad cate gag gag gag gag gat and gac gag gag gat gag gag gag gag gag gag gag	gry of the ser ser ser ser ser ser ser ser ser se	ggg gga cat tca ttg aaa ggg cag ata gga ggg cat tca ttg tca ttg gla aaa aaa aca cag gaa gga gga ggc cat tta atc gaa ggg caa ttg gla gaa ggg caa aaa aaa aca ttg tgt gaa gga ggg cat tta atc gaa ggg caa ttg gla gaa ggg caa aaa aaa aca ttg gat gaa ggg caa ata gaa ggg caa ttg tca ttg that atc gaa ggg ggc caa ttg gaa gga ggg caa aaa aaa acc gga gga ttg tca gga ata ggg gaa ggg caa ttg gaa ggg aaa aaa aaa acc gga gaa ggg aaa aaa	ggg gga cat tca ttg aaa aac ggt ggt gtg gtg ata gac gag aac ttc cgg ggg gga cat tca ttg aac gat gac gtg gtg ttg ttt tcg gtg gac ggt gtg ttg ttt tcg gac atc gat ggc gtg gtg ata ggc ggc ggc atc ggc Arg 1005  ggg gga cat tca ttg aaa ggc ggt gtg ata Mar Ala Val Ala 1015  aaa tcg ggt gtg ctc gga atc ggt gtg 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Lys Asp Val Leu Glu Lys Ser Met Asn Val Ile Ile Gly Arg Tyr Asp 50 60	
Ile Phe Arg Thr Val Phe Val His Glu Lys Met Lys Arg Pro Val Gln 65 70 75 80	

Val Val Leu Lys Glu Arg Ser Phe Gln Ala Glu Glu Ile Asp Leu Ser 85 90 95

Gly Leu Ser Glu Ala Glu Gln Asn Glu Arg Ile Glu Asp Tyr Lys Arg 100 105 110

Lys Asp Lys Glu Lys Gly Phe Asn Leu Ser Lys Asp Ile Pro Met Arg 115 120 125 Page 73

Thr Ala Val Phe Lys Lys Gly Gln Asp Arg Tyr Glu Trp Val Trp Ser 130 135 140 Tyr His His Ile Leu Leu Asp Gly Trp Cys Phe Gly Ile Val Val Gln 145 150 150 160 Glu Leu Phe Glu Val Tyr Asn Ala Leu Arg Glu Asn Arg Pro Tyr Ser 165 170 175 Leu Gly Pro Val Arg Pro Tyr Lys Glu Tyr Ile Lys Trp Leu Glu Ser 180 185 190 Gln Asp Lys Gln Lys Ser Leu Val Tyr Trp Glu His His Leu Ala Gly 195 200 205 Phe Glu Gly Gln Thr Thr Phe Ser Glu Gln Arg Lys Asn Pro Glu Gln 210 215 220 Ala Gly Asn Gln Pro Glu Glu Leu Leu Phe Thr Leu Pro Lys Glu Asp 225 230 235 240 Thr Glu Ala Phe Thr Gln Leu Ala Lys Ala His His Thr Thr Leu Ser 245 250 255 Thr Ala Leu Gln Ala Val Trp Ser Val Leu Leu Ser Arg Tyr Gln Arg 260 265 270 Ser Arg Asp Leu Ile Phe Gly Thr Val Val Ser Gly Arg Pro Ala Asp 275 280 285 Ile Gln Gly Val Glu His Met Val Gly Leu Phe Ile Asn Val Val Pro 290 295 300 Lys Arg Ala Thr Phe Gly Pro Gln Thr Thr Phe Thr Glu Leu Val Ala 305 310 315 320 Asp Leu Gln Lys Gln Ala Leu Glu Ala Glu Pro His Gln Tyr Val Pro 325 330 335 Leu Tyr Glu Ile Gln Ser His Ile Ala Ala Pro Asn Leu Ile Asp His 340 350 Ile Ile Val Phe Glu Asn Tyr Pro Leu Gln Glu Ala Asn Lys Gln Gln 355 360 365 Lys Glu Lys Asn Leu Gly Phe Thr Met Asp Glu Ile Ser Val Phe Glu 370 380 Lys Ser Asn Tyr Asp Leu Asn Leu Leu Ala Ser Pro Gly Glu Glu Met 385 390 395 400 Page 74

Met Leu Lys Leu Ala Tyr Asn Arg Asn Ile Phe Glu Pro Ser Phe Ile 405 415 Leu Arg Leu Lys Glu Gln Leu Leu Thr Ala Ile Arg Glu Val Ile Arg 420 425 430 Asp Pro Glu Gln Pro Leu Asp Asp Ile Cys Val Val Ser Lys Lys Glu 435 440 445 Lys Glu Arg Leu Leu His Glu Phe Asn Gly Gln Val Gln Arg Lys Gln 450 460 Ile Gly Leu Thr Ile Pro Gln Trp Phe Glu Gln Tyr Ala Glu Ala Leu 465 470 475 480 Pro Glu Lys Pro Ala Val Ser Ala Ala Gly Arg Thr Leu Thr Tyr Arg 485 490 495 Glu Leu Asn Glu Lys Ala Asn Lys Val Ala His Leu Leu Arg Lys Lys 500 505 510 Asn Val Gly Arg Gly Glu Pro Val Ala Leu Leu Phe Arg Arg Ser Pro 515 520 525 Glu Met Val Ile Ala Ile Leu Ala Val Leu Lys Ala Gly Gly Ala Tyr 530 535 540 Leu Pro Ile Asp Pro Glu Tyr Pro Glu Ala Arg Ile Gln Tyr Met Leu 545 550 555 560 Glu Asp Ser Gly Ala Val Cys Met Leu Thr Gln Glu Glu Leu Ala Gly 565 575 Arg Ala Ala Ser Leu Ser Phe His Lys His Thr Ile Leu Ile Asp Asp 580 585 590 Pro Ala Val Ser Ala Glu Ser Gly Arg Asn Leu Glu Ile Ala Ala Glu 595 600 605 Pro Asp Asp Leu Ala Tyr Ile Met Tyr Thr Ser Gly Thr Thr Gly Lys 610 615 620 Pro Lys Gly Asn Leu Thr Thr His Ala Asn Ile Thr Arg Val Val Lys 625 630 635 640 Glu Thr Asn Tyr Ile Ser Leu Ser Glu Lys Asp Thr Leu Leu Ser Leu 645 650 655 Ser Asn Tyr Ala Phe Asp Gly Phe Thr Phe Asp Leu Tyr Gly Ala Leu 660 665 670 Page 75

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- Val Ala Asp Gly Ile Ala Lys Lys Met Thr Ala Tyr Tyr Ser Tyr 1190 1195 1200 Page 77

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Page 78	

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His Leu Trp Thr Met Lys Glu Ser Phe Ile Lys Gln Ala Gly Lys Gly 145 150 155 160

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Arg Ala Ser Val Glu Thr Pro Pro Gly Tyr Pro Pro Cys Tyr Ile Gln 180 185 190

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DNA Bacillus licheniformis

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